

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/801,486A
Source: JFW/6
Date Processed by STIC: 9/28/06

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IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/801,486A

DATE: 09/28/2006

TIME: 08:50:34

Input Set : A:\00281bus.txt
 Output Set: N:\CRF4\09282006\J801486A.raw

4 <110> APPLICANT: Yan, Riqiang
 5 Tomasselli, Alfredo G.
 6 Gurney, Mark E.
 7 Emmons, Thomas L.
 8 Bienkowski, Mike J.
 9 Heinrikson, Robert L.
 11 <120> TITLE OF INVENTION: SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY
 13 <130> FILE REFERENCE: 29915/00281BUS
 15 <140> CURRENT APPLICATION NUMBER: 10/801,486A
 16 <141> CURRENT FILING DATE: 2004-03-16
 18 <150> PRIOR APPLICATION NUMBER: 09/908,943
 19 <151> PRIOR FILING DATE: 2001-07-19
 21 <150> PRIOR APPLICATION NUMBER: 60/219,795
 22 <151> PRIOR FILING DATE: 2000-07-19
 24 <160> NUMBER OF SEQ ID NOS: 199
 26 <170> SOFTWARE: PatentIn Ver. 2.0
 28 <210> SEQ ID NO: 1
 29 <211> LENGTH: 2070
 30 <212> TYPE: DNA
 31 <213> ORGANISM: Homo sapiens
 33 <400> SEQUENCE: 1
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 35 ggcacccagc acggcatccg gctgcccctg cgcagccgccc tggggggcgc cccctgggg 120
 36 ctgcggctgc cccgggagac cgacgaagag cccgaggagc cccggccggag gggcagctt 180
 37 gtggagatgg tggacaacct gagggcaag tcggggcagg gctactacgt ggagatgacc 240
 38 gtgggcagcc ccccgccagac gctcaacatc ctgggtggata caggcagcag taactttgca 300
 39 gtgggtgctg ccccccaccc ctgcctgcat cgctactacc agaggcagct gtccagcaca 360
 40 taccgggacc tccggaaggg tggatgtgtg ccctacaccc agggcaagt ggaaggggag 420
 41 ctgggcaccc acctggtaag catccccat ggcaccaacg tcactgtgcg tgccaaacatt 480
 42 gctgccatca ctgaatcaga caagttcttc atcaacggct ccaactggga aggcatcctg 540
 43 gggctggcct atgctgagat tggcaggcct gacgactccc tggagcctt ctttactct 600
 44 ctggtaaaggc agacccacgt tcccaaccc ttctccctgc acctttgtgg tgctggctc 660
 45 cccctcaacc agtctgaagt gctggcctct gtcggaggga gcatgatcat tggaggtatc 720
 46 gaccactcgc tgtacacagg cagtctctgg tatacaccca tccggccggga gtggattat 780
 47 gaggtcatca ttgtgggggt ggagatcaat ggacaggatc tgaaaatgga ctgcaaggag 840
 48 tacaactatg acaagagcat tggagacagt ggcacccacca accttcgtt gcccaagaaa 900
 49 gtgtttgaag ctgcagtc当地 atccatcaag gcagctctt ccacggagaa gttccctgtat 960
 50 ggtttctggc taggagagca gctgggtgtgc tggcaagcag gcaccacccc ttggAACATT 1020
 51 ttcccagtca tctcaactca cctaatgggt gaggttacca accagtccctt ccgcacatcacc 1080
 52 atccctccgc agcaataacct gcccgcagtg gaagatgtgg ccacgtccca agacgactgt 1140
 53 tacaagtttgc ccatctcaca gtcatccacg ggcactgtta tggagactgt tatcatggag 1200
 54 ggcttctacg ttgtcttga tcggggcccgaa aaacgaattt gctttgtgtt cagcgcttgc 1260
 55 catgtgcacg atgagttcag gacggcagcg gttggaaaggcc cttttgtcac cttggacatg 1320

(CPG. 6-7)

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56 gaagactgtg gctacaacat tccacagaca gatgagtcaa ccctcatgac catagcctat 1380
 57 gtcatggctg ccatctgcgc cctcttcatg ctgccactct gcctcatgg gtgtcagtgg 1440
 58 cgctgcctcc gctgcctgctg ccagcagcat gatgactttg ctgatgacat ctccctgctg 1500
 59 aagtggagg gcccattggc agaaagataga gattcccctg gaccacacct ccgtggttca 1560
 60 ctttggtcac aagttaggaga cacagatggc acctgtggcc agagcacctc aggaccctcc 1620
 61 ccacccacca aatgcctctg ctttgcatttga gaaggaaaag gctggcaagg tgggttccag 1680
 62 ggactgtacc tggtaggaaac agaaaagaga agaaaagaagc actctgtctgg cgggaataact 1740
 63 cttggtcacc tcaaatttaa gtcgggaaat tctgctgctt gaaacttcag ccctgaacct 1800
 64 ttgtccacca ttccctttaaa ttctccaacc caaagtattt ttcttttctt agtttcagaa 1860
 65 gtactggcat cacacgcagg ttaccttggc gtgtgtccct gtggtaggat ggcagagaag 1920
 66 agaccaagct tgtttcctg ctggccaaag tcagtaggag agatgcaca gtttgctatt 1980
 67 tgcttttagag acagggactg tataaaacaag cctaacattt gtcggcaagat tgcctcttga 2040
 68 attaaaaaaaaaaaaaaa 2070

70 <210> SEQ ID NO: 2

71 <211> LENGTH: 501

72 <212> TYPE: PRT

73 <213> ORGANISM: Homo sapiens

75 <400> SEQUENCE: 2

76 Met Ala Gln Ala Leu Pro Trp Leu Leu Trp Met Gly Ala Gly Val
 77 1 5 10 15
 79 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
 80 20 25 30
 82 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
 83 35 40 45
 85 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
 86 50 55 60
 88 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
 89 65 70 75 80
 91 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
 92 85 90 95
 94 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
 95 100 105 110
 97 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
 98 115 120 125
 100 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
 101 130 135 140
 103 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
 104 145 150 155 160
 106 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
 107 165 170 175
 109 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp
 110 180 185 190
 112 Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro
 113 195 200 205
 115 Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
 116 210 215 220
 118 Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
 119 225 230 235 240
 121 Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg

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122	245	250	255
124	Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln		
125	260	265	270
127	Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val		
128	275	280	285
130	Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala		
131	290	295	300
133	Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp		
134	305	310	315
136	Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr		
137	325	330	335
139	Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val		
140	340	345	350
142	Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg		
143	355	360	365
145	Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala		
146	370	375	380
148	Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu		
149	385	390	395
151	Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala		
152	405	410	415
154	Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu		
155	420	425	430
157	Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro		
158	435	440	445
160	Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala		
161	450	455	460
163	Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp		
164	465	470	475
166	Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp		
167	485	490	495
169	Ile Ser Leu Leu Lys		
170	500		
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174	<211> LENGTH: 1977		
175	<212> TYPE: DNA		
176	<213> ORGANISM: Homo sapiens		
178	<400> SEQUENCE: 3		
179	atggcccaag ccctgccctg gtcctgtcg tggatggcg cgggagtgt gcctgcccac 60		
180	ggcacccagc acggcatccg gctgccctg cgcagccgac tggggggcgc cccctgggg 120		
181	ctgcggctgc cccggagac cgacgaagag cccgaggagcc cccggccggag gggcagctt 180		
182	gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240		
183	gtgggcagcc ccccgagac gctcaacatc ctggggata caggcagcag taactttgca 300		
184	gtgggtgctg ccccccaccc ttccctgtcat cgctactacc agaggcagct gtccagcaca 360		
185	taccgggacc tccgaaagggtgttatgtg ccctacaccc agggcaagtg ggaaggggag 420		
186	ctgggcaccc acctggtaag catccccat ggcccaacg tcactgtgcg tgccaaacatt 480		
187	gtgccatca ctgaatcaga caagttcttc atcaacggct ccaactggga aggcacccctg 540		
188	gggctggcct atgctgagat tgccaggctt tgtgtgtcg gcttccccct caaccagtct 600		
189	gaagtgctgg cctctgtcgg agggagcatg atcattggag gtatcgacca ctcgctgtac 660		

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190 acaggcagtc tctggatac acccatccgg cgggagtgg attatgaggt gatcattgtg 720
 191 cgggtggaga tcaatggaca gatctgaaa atggactgca aggagtacaa ctatgacaag 780
 192 agcattgtgg acagttggcac caccacccctt cggttgcacca agaaagtgtt tgaagctgca 840
 193 gtcaaattcca tcaaggcagc ctccctccacg gagaagttcc ctgatggttt ctggcttagga 900
 194 gagcagctgg tgtgctggca agcaggcacc accccttggaa acatttccc agtcatctca 960
 195 ctctacctaa tgggtgaggt taccacccag tccttcgcac tcaccatctt tccgcagcaa 1020
 196 tacctgcggc cagtggaa tggccacg tcccaagacg actgttacaa gtttgccatc 1080
 197 tcacagtcat ccacgggcac tggttatggaa gctgttatca tggagggctt ctacgttgc 1140
 198 ttgtatcgaa cccgaaaacg aattggctt gctgtcagcg cttgcccattgt gcacgatgag 1200
 199 ttcaggacgg cagcgggtggaa aggccctttt gtcaccccttgg acatggaa gatggctac 1260
 200 aacattccac agacagatga gtcaaccctc atgaccatag cctatgtcat ggctgcccattc 1320
 201 tgcccccctct tcattgtgcc actctgcctc atgggtgtc agtggcgcctg cttccgcgtc 1380
 202 ctgcgcctcaggc agcatgtatga ctttgcgtat gacatctccc tgctgaagtg aggaggccca 1440
 203 tgggcagaag atagagattc ccctggacca cacccctgt gttcactttg gtcacaagta 1500
 204 ggagacacag atggcacctg tggccagagc acctcaggac cctcccccacc caccataatgc 1560
 205 ctctgccttg atggagaagg aaaaggctgg caaggtgggt tccaggact gtacctgttag 1620
 206 gaaacagaaa agagaagaaa gaagcactt gctggcggga atactctgg tcacctcaaa 1680
 207 tttaagtctgg gaaattctgc tgcttgaaac ttccatccctg aacctttgtc caccattctt 1740
 208 tttaattcttc caaccctaaag tattttttt ttcttagttt cagaagtact ggcacacac 1800
 209 gcaggttacc ttggcgtgtg tccctgtggt accctggcag agaagagacc aagttgttt 1860
 210 ccctgtggc caaagtctgttggaggat gcacagtttgcctttagagacagg 1920
 211 gactgtataa acaaggctaa cattggtgca aagattgcctt tgaaaaaaa aaaaaaaa 1977

213 <210> SEQ ID NO: 4

214 <211> LENGTH: 476

215 <212> TYPE: PRT

216 <213> ORGANISM: Homo sapiens

218 <400> SEQUENCE: 4

219 Met Ala Gln Ala Leu Pro Trp Leu Leu Trp Met Gly Ala Gly Val
 220 1 5 10 15
 222 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
 223 20 25 30
 225 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
 226 35 40 45
 228 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
 229 50 55 60
 231 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
 232 65 70 75 80
 234 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
 235 85 90 95
 237 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
 238 100 105 110
 240 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
 241 115 120 125
 243 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
 244 130 135 140
 246 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
 247 145 150 155 160
 249 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
 250 165 170 175

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252 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Leu Cys Gly
 253 180 185 190
 255 Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
 256 195 200 205
 258 Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
 259 210 215 220
 261 Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val
 262 225 230 235 240
 264 Arg Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr
 265 245 250 255
 267 Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu
 268 260 265 270
 270 Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser
 271 275 280 285
 273 Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val
 274 290 295 300
 276 Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser
 277 305 310 315 320
 279 Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile
 280 325 330 335
 283 Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln
 284 340 345 350
 286 Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val
 287 355 360 365
 289 Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala
 290 370 375 380
 292 Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu
 293 385 390 395 400
 295 Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu
 296 405 410 415
 298 Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr
 299 420 425 430
 301 Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu
 302 435 440 445
 304 Cys Leu Met Val Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln
 305 450 455 460
 307 His Asp Asp Phe Ala Asp Asp Ile Ser Leu Leu Lys
 308 465 470 475
 311 <210> SEQ ID NO: 5
 312 <211> LENGTH: 14
 313 <212> TYPE: PRT
 314 <213> ORGANISM: Artificial Sequence
 316 <220> FEATURE:
 317 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
 318 peptide sequence
 320 <400> SEQUENCE: 5
 321 Lys Val Glu Ala Asn Tyr Glu Val Glu Gly Glu Arg Lys Lys
 322 1 5 10
 325 <210> SEQ ID NO: 6

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; Xaa Pos. 7
Seq#:15; Xaa Pos. 4,7
Seq#:16; Xaa Pos. 1,4,5,6,7
Seq#:17; Xaa Pos. 1,2,4,5,6,7
Seq#:18; Xaa Pos. 1,2,4,5,6,7
Seq#:21; Xaa Pos. 5
Seq#:27; Xaa Pos. 7,19
Seq#:28; Xaa Pos. 6,7,11,20
Seq#:41; Xaa Pos. 9
Seq#:49; Xaa Pos. 1
Seq#:50; Xaa Pos. 2
Seq#:51; Xaa Pos. 3
Seq#:52; Xaa Pos. 4
Seq#:53; Xaa Pos. 5
Seq#:54; Xaa Pos. 6
Seq#:55; Xaa Pos. 7
Seq#:56; Xaa Pos. 8
Seq#:57; Xaa Pos. 1
Seq#:58; Xaa Pos. 2
Seq#:59; Xaa Pos. 3
Seq#:60; Xaa Pos. 4
Seq#:61; Xaa Pos. 5
Seq#:62; Xaa Pos. 6
Seq#:63; Xaa Pos. 7
Seq#:64; Xaa Pos. 8
Seq#:65; Xaa Pos. 1
Seq#:66; Xaa Pos. 2
Seq#:67; Xaa Pos. 3
Seq#:68; Xaa Pos. 4
Seq#:69; Xaa Pos. 5
Seq#:70; Xaa Pos. 6
Seq#:71; Xaa Pos. 7
Seq#:72; Xaa Pos. 8
Seq#:73; Xaa Pos. 1
Seq#:74; Xaa Pos. 2
Seq#:75; Xaa Pos. 3
Seq#:76; Xaa Pos. 4
Seq#:77; Xaa Pos. 7
Seq#:78; Xaa Pos. 8
Seq#:79; Xaa Pos. 8
Seq#:80; Xaa Pos. 9
Seq#:81; Xaa Pos. 1,7
Seq#:82; Xaa Pos. 2,7
Seq#:83; Xaa Pos. 3,7

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Seq#:84; Xaa Pos. 4,7
Seq#:85; Xaa Pos. 5,7
Seq#:86; Xaa Pos. 6,7
Seq#:87; Xaa Pos. 7
Seq#:88; Xaa Pos. 7,8
Seq#:89; Xaa Pos. 1
Seq#:90; Xaa Pos. 1,2

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L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
L:476 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:0
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:595 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
M:341 Repeated in SeqNo=27
L:731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
L:1045 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:0
L:1064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0
L:1083 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:0
L:1102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:0
L:1121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0
L:1140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:0
L:1159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:0
L:1178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:0
L:1197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:0
L:1216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
L:1235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60 after pos.:0
L:1273 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
L:1292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0
L:1311 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:0
L:1330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
L:1349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
L:1368 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
L:1387 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67 after pos.:0
L:1406 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:0
L:1426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:0
L:1445 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0
L:1464 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:0
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L:1502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:0
L:1521 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0
L:1540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75 after pos.:0
L:1559 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76 after pos.:0
L:1578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:0
L:1597 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:0
L:1616 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 after pos.:0
L:1635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 after pos.:0
L:1659 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:0
L:1683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:0
L:1707 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84 after pos.:0
L:1755 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:0

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L:1779 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:0

L:1798 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87 after pos.:0